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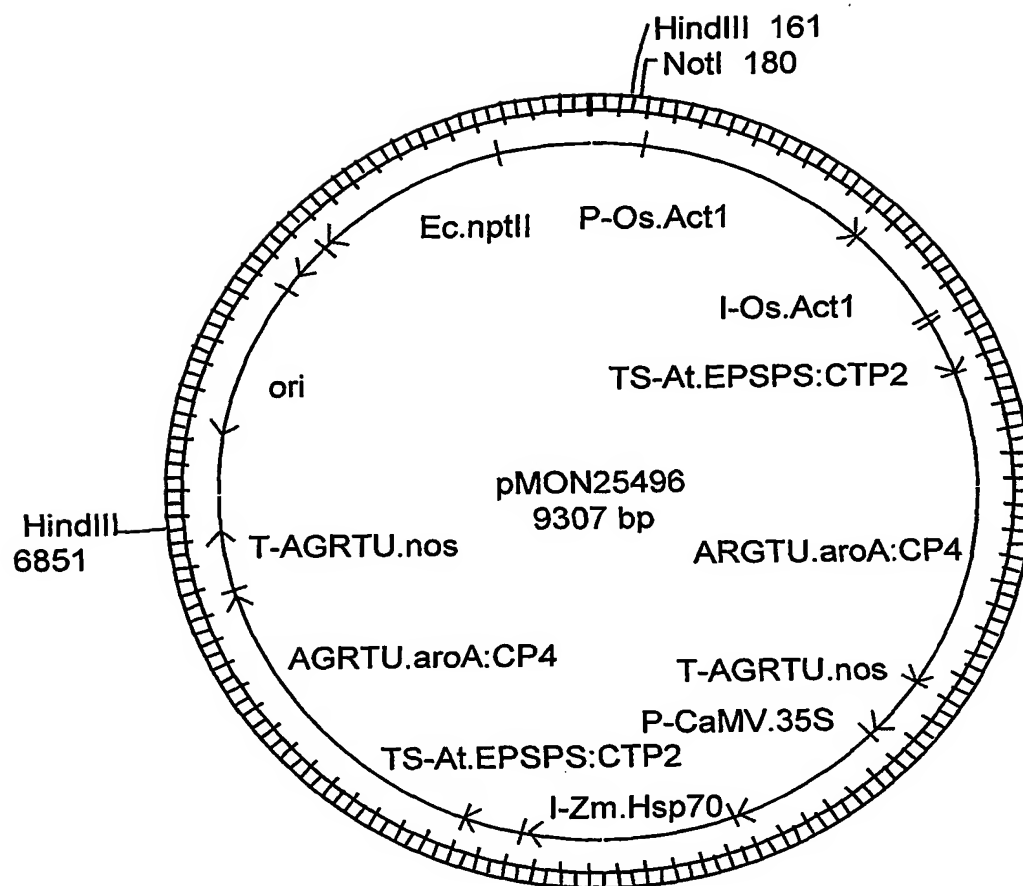


Figure 1

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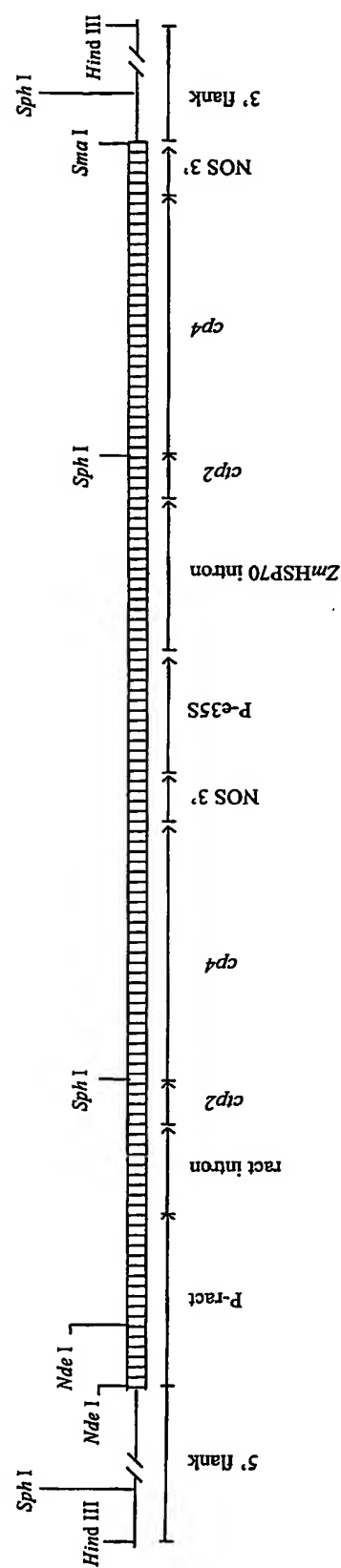


Figure 2

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1 aagcgagtat cctgATAAGA AAGGAAGAAG ACGATCGCTC TGTCTATGGG
51 CGGGGCTCAG GGCGACGACA GAACCAGAGC TTTCGTCGTG AACAAAACAG
101 GGAAGGACCA AAGCAGAGGA AGAGGAGAGG AAACAGAGAG AAAGAGGGGG
151 TTGGTAGGTA CTTGGTGGTC CCTGCTACTT CTCCAACAGC AGCAGAAAGG
201 AAAGAAGAAC GAACCAAGGC ACAAGTACGC TCCAACCGAG CCATCCCTTT
251 CTTCCCTTTA TCATTGACTT TAATCATGAG AAATCTAATT AATTAATTAA
301 ACTCTACGCA AAAGGCATAT AAAATTGTCA ATTATGCAAG GCAGTTGCCC
351 TGTTTCTGGT AGCCGGTTAC AACACAGGAA GACAACCAA AGCGTCGGAA
401 AAGTGAGTTT AGTCGAATCT GAATTCAATG TGAAAGATTT TTGTAAAGAA
451 TGAAATAAAT CCCGATAAAA AAAGAATGAA CAAAAGGAAA CTAAAAAACT
501 GTGGATGTGA GTCCAACGTT TAAGCATATC GATGCAAACG TGATGAAGAA
551 CCAAACGCGC CGGCGGAAGA CGGATTCCCG GAAGACCAA TTAAAGACGA
601 TAGTTGTCGA GCAAACGACC AAAAGAAGAA GATCCGACAT ATGCTTAAGA
651 AGAGAGTCGG GATAGTCCAA AATAAAACAA AGGTAAGATT ACCTGGTCAA
701 AAGTGAAAAC ATCAGTTAAA AGGTGGTATA AAGTAAAATA TCGGTAATAA
751 AAGGTGGCCC AAAGTGAAAT TTACTCTTTT CTACTATTAT AAAAATTGAG
801 GATGTTTTTG TCGGTACTTT GATACGTCAT TTTTGTATGA ATTGGTTTTT
851 AAGTTTATTC GCTTTTGGAA ATGCATATCT GTATTGagt cggqgtt

Figure 3

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1 agattgaatc ctGTTGCCGG TCTTGCGATG ATTATCATAT AATTTCTGTT
51 GAATTACGTT AAGCATGTAA TAATTAACAT GTAATGCATG ACGTTATTTA
101 TGAGATGGGT TTTTATGATT AGAGTCCCGC AATTATACAT TTAATACGCG
151 ATAGAAAACA AAATATAGCG CGCAAAC TAG GATAAATTAT CGCGCGCGGT
201 GTCATCTATG TTAGTAGATC GGGGATATCC CCGGGGAATT CGGTACCATG
251 TACCACGGAA CAGAAAAAAG AAAGGCCAC GGTGTGCAG GAAACGGCCA
301 CCGCGCGAGC CAGCGCCTCA CGCCTCATCC GCCATTCCGT CGAGCACCCC
351 GCACGCGCCG CCGCTGCTAT GCTCCTCCGG CCGCGCCCCT TCCTCCTCCA
401 GGTCTTCAGC CCGCTTCGCT CCTCCGCGC CCCCCTCGCG GTCCGCCGCA
451 CGCTCTCAGC gcacgccgcg qcaq

Figure 4

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5' GACATATGCTTAAGAAGAGAGTCG 3' (SEQ ID NO:1)
5' AATTCGGTACCATGTACCACGAAC 3' (SEQ ID NO:2)

Figure 5